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PLEASE DELIVER TO ATTN: Examiner Seharaseyon, J.

Concerning the following applications:

ATTORNEY DOCKET NO.: 9409/2023D

Application of:	Samson, et al.	Examiner:	Seharaseyon, J.
Serial No.:	09/938,703	Group Art Unit:	1647
Filed:	August 24, 2001		
Entitled:	"CCR5 Chemokine Receptor-Specific Antibodies"	Conf. No.:	1978

ATTORNEY DOCKET NO.: 9409/2023B

Application of:	Samson, et al.	Examiner:	Seharaseyon, J.
Serial No.:	09/938,719	Group Art Unit:	1647
Filed:	August 24, 2001		
Entitled:	"Nucleic Acids Encoding Active and Inactive CCR5 Chemokine Receptors"	Conf. No.:	3864

Facsimile No.: 1-703-746-5177

Number of pages including cover sheet: 6

Transmitted herewith: Supplemental Preliminary Amendment

CERTIFICATION

I hereby certify that the paper identified above is being facsimile transmitted to the U.S. Patent and Trademark Office, on the date indicated below.

Date: June 5 , 2003


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Attorney Docket N .: 9409/2023D PATENT

THE UNITED STATES PATENT AND TRADEMARK OFFICE

Application of:	Samson, et al.	Examiner:	Seharaseyon, J.
U.S. Serial No.:	09/938,703	Group Art Unit:	1647
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(AF)

Commissioner for Patent
P.O. Box 1450
Alexandria, VA 22313-1450

SUPPLEMENTAL PRELIMINARY AMENDMENTIn the Claims

Please replace claims 67, 68, 72, and 73 with the following claims. A marked up version of the claim amendments is filed herewith.

F1 50
67. An antibody which binds to a polypeptide comprising amino acid residues 185 to 215 of SEQ ID NO: 6.

F2 52
68. An antibody which binds to a polypeptide consisting of amino acid residues 185 to 215 of SEQ ID NO: 6.

F3 61
72. An antibody which binds to a polypeptide comprising at least four contiguous amino acid residues of the sequence of residues 185 to 215 of SEQ ID NO: 6.

? 4 62
73. An antibody which binds to a polypeptide consisting at least four contiguous amino acid residues of the sequence of residues 185 to 215 of SEQ ID NO: 6.

REMARKS

Upon entry of this amendment, claims 67-76 are pending. No new matter is introduced by this amendment. Support for the newly added claims may be found throughout the specification and the sequence listing.